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		STATES OF	Washington, D.C. 202	231	ti
APPLICATION NO.	FILING DATE	FIRST NAMED I	NVENTOR	A.	TTORNEY DOCKET NO.
09/234,182	01/20/9	99 HSEI	,	V	P1085R4-1A
			7	E	XAMINER
RICHARD B	L CALLES	HM12/0322	r.	IBRIN	rīu tvi
GENENTECH				<u>UNIT</u>	PAPER NUMBER
1 DNA WAY					
SOUTH SAN	FRANCISCO	CA 94080-4990	1 :	544	/
			DATE MA	AILED:	
					03/22/00

Please find below and/or attached an Office communication concerning this application or proceeding.

Commissioner of Patents and Trademarks



UNITED STATES DEPARTMENT OF COMMERCE Patent and Trademark Office
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SERIAL NUMBER	FILING DATE	FIRST NAMED APPLICANT	ATTORNEY DOCKET NO.

EXAMINER

ART UNIT PAPER NUMBER

DATE MAILED:

Please find below a communication from the EXAMINER in charge of this application

This application contains sequence disclosures that are encompassed by the definitions for nucleotide and/or amino acid sequences set forth in 37 CFR 1.821(a)(1) and (a)(2). However, this application fails to comply with the requirements of 37 CFR 1.821 through 1.825 for the reason(s) set forth on the attached Notice To Comply With Requirements For Patent Applications Containing Nucleotide Sequence And/Or Amino Acid Sequence Disclosures. See the attached copy of the marked-up "Raw Sequence Listing."

Any inquiry concerning this communication should be directed to Examiner **Marianne DiBrino**, Art Unit **1644**, whose telephone number is **703-305-1987**.

Any inquiry of a general nature or relating to the status of this application should be directed to the Technology Center receptionist whose telephone number is (703) 308-0196.

APPLICANT IS GIVEN A ONE MONTH EXTENDABLE PERIOD WITHIN WHICH TO COMPLY WITH THE SEQUENCE RULES, 37 CFR 1.821 - 1.825. Failure to comply with these requirements will result in ABANDONMENT of the application under 37 CFR 1.821(g). Extensions of time may be obtained by filing a petition accompanied by the extension fee under the provisions of 37 CFR 1.136. In no case may an applicant extend the period for response beyond the six month statutory period. Applicant is requested to return a copy of the attached Notice to Comply with the response.

Marianne DiBrino, Ph.D.

Mr. ringare

March 19, 2000

CHRISTINA Y. CHAN
SUPERVISORY PATENT EXAMINER
GROUP 1800

Application No.: 09/234/8

The nucleotide and/or amino acid sequence disclosure contained in this application does not comply with the requirements for such a disclosure as set forth in 37 C.F.R. 1.821 - 1.825 for the following reason(s)
1. This application deady falls to comply with the requirements of 37 C.F.R. 1.821-1.825. Applicant's attention is directed to these regulations, published at 1114 OG 29, May 15, 1990 and at 55 FR 18230, May 1, 1990.
2. This application does not contain, as a separate part of the disclosure on paper copy, a "Sequence Usting" as required by 37 C.F.R. 1.821(c).
3. A copy of the "Sequence Listing" in computer readable form has not been submitted as required by 37 C.F.R. 1.821(e).
A. A copy of the "Sequence Listing" in computer readable form has been submitted. However, the content of the computer readable form does not comply with the requirements of 37 C.F.R. 1.822 and/or 1.823, as indicated on the attached copy of the marked -up "Raw Sequence Listing."
5. The computer readable form that has been filed with this application has been found to be damaged and/or unreadable as indicated on the attached CRF Diskette Problem Report. A Substitute computer readable form must be submitted as required by 37 C.F.R. 1.825(d).
6. The paper copy of the "Sequence Listing" is not the same as the computer readable from of the "Sequence Listing" as required by 37 C.F.R. 1.821(e).
7. Other:
Applicant Must Provide:
An lattitude computer readable form (CRF) copy of the "Sequence Listing".
An lattick or substitute paper copy of the "Sequence Listing", as well as an amendment directing its entry into the specification.
A statement that the content of the paper and computer readable copies are the same and, where

applicable, include no new matter, as required by 37 C.F.R. 1.821(e) or 1.821(f) or 1.821(g) or 1.825(b) or 1.825(d).

For questions regarding compliance to these requirements, please contact:

For Rules Interpretation, call (703) 308-4216

For CRF Submission Help, call (703) 308-4212

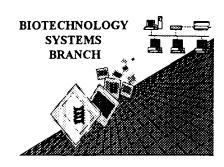
For Patentin software help, call (703) 308-6856

PLEASE RETURN A COPY OF THIS NOTICE WITH YOUR RESPONSE

COPY FOR [] File [] Applicant

Dibrino

RAW SEQUENCE LISTING ERROR REPORT



The Biotechnology Systems Branch of the Scientific and Technical Information Center (STIC) detected errors when processing the following CRF diskette:

Application Serial Number:

09/234,182

Art Unit / Team No.:

1600

Date Processed by STIC:

1/29/99

THE ATTACHED PRINTOUT EXPLAINS THE ERRORS DETECTED.

PLEASE BE SURE TO FORWARD THIS INFORMATION TO THE APPLICANTS BY EITHER:

- 1) INCLUDING A COPY OF THIS PRINTOUT IN YOUR NEXT COMMUNICATION TO THE APPLICANTS ALONG WITH A NOTICE TO COMPLY or,
- 2) CALLING APPLICANTS AND FAXING THEM A COPY OF THE PRINTOUT WITH A NOTICE TO COMPLY

THIS WILL INSURE THAT THE NEXT SUBMISSION RECEIVED FROM THEM WILL BE ERROR FREE.

IF YOU HAVE ANY FURTHER QUESTIONS, PLEASE CALL:

MARK SPENCER 703-308-4212

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PAGE: 1

RAW SEQUENCE LISTING PATENT APPLICATION US/09/234,182

DATE: 01/29/1999 TIME: 13:42:11

INPUT SET: S30387.raw

This Raw Listing contains the General Information Section and those Sequences containing ERRORS.

1		sequence Listing Not Comply
2		Constal Information: Corre
3	(1)	General Information:
4		
5	(i)	APPLICANT: Genentech, Inc., Hsei, Vanessa
6		Koumenis, Iphigenia
7 8		Leong, Steven R. Presta, Leonard G.
9		Shahrokh, Zahra
10		Zapata, Gerardo A.
11		Dapaca, Octatao A.
12	(ii)	TITLE OF INVENTION: ANTIBODY FRAGMENT-POLYMER CONJUGATES AND
13	, ,	HUMANIZED ANTI-IL-8 MONOCLONAL ANTIBODIES
14		
15	(iii)	NUMBER OF SEQUENCES: 72
16		
17	(iv)	CORRESPONDENCE ADDRESS:
18		(A) ADDRESSEE: Genentech, Inc.
19		(B) STREET: 1 DNA Way
20		(C) CITY: South San Francisco
21 22		(D) STATE: California (E) COUNTRY: USA
23		(E) COUNTRY: USA (F) ZIP: 94080
24		(r) Elr. 94000
25	(v)	COMPUTER READABLE FORM:
26	, , ,	(A) MEDIUM TYPE: 3.5 inch, 1.44 Mb floppy disk
27		(B) COMPUTER: IBM PC compatible
28		(C) OPERATING SYSTEM: PC-DOS/MS-DOS
29		(D) SOFTWARE: WinPatin (Genentech)
30		
31	(vi)	CURRENT APPLICATION DATA:
32		(A) APPLICATION NUMBER:
33		(B) FILING DATE: 20-Jan-1999
34 35		(C) CLASSIFICATION:
36	(wii)	PRIOR APPLICATION DATA:
37	(• ± ± /	(A) APPLICATION NUMBER: 60/074330
38		(B) FILING DATE: 22-JAN-1998
39		(2) (2223)
40	(vii)	PRIOR APPLICATION DATA:
41		(A) APPLICATION NUMBER: 60/094003
42		(B) FILING DATE: 24-JUL-1998
43		
44	(vii)	PRIOR APPLICATION DATA:
45		(A) APPLICATION NUMBER: 60/094013

PAGE: 1

SEQUENCE VERIFICATION REPORT PATENT APPLICATION US/09/234,182

DATE: 01/29/1999 TIME: 13:42:12

INPUT SET: S30387.raw

Line	Error	Original Text
413 1155 1254	Wrong or Missing Sequence Type Wrong or Missing Sequence Type Wrong or Missing Sequence Type	(B) TYPE: DNA (B) TYPE: PRT (B) TYPE: PRT

PAGE: 2

RAW SEQUENCE LISTING PATENT APPLICATION US/09/234,182

DATE: 01/29/1999 TIME: 13:42:11

INPUT SET: S30387.raw

46		(B) FILING DATE: 24-JUL-1998
47		
48	(vii)	PRIOR APPLICATION DATA:
49		(A) APPLICATION NUMBER: 60/075467
50		(B) FILING DATE: 20-FEB-1998
51		
52	(viii)	ATTORNEY/AGENT INFORMATION:
53		(A) NAME: Love, Richard B.
54		(B) REGISTRATION NUMBER: 34,659
55		(C) REFERENCE/DOCKET NUMBER: P1085R4-1A
56		
57	(ix)	TELECOMMUNICATION INFORMATION:
58		(A) TELEPHONE: 650/225-5530
59		(B) TELEFAX: 650/952-9881

ERRORED SEQUENCES FOLLOW:

	409	(2) INFORMATION FOR SEQ ID NO:22:
>	410 411 412 413 414 415 416	(i) SEQUENCE CHARACTERISTICS: (A) LENGTH: 21 base pairs (B) TYPE: DNA hurlue and (ONA is only used with here (C) STRANDEDNESS: Single (D) TOPOLOGY: Linear Rule from
	417 418 419	(xi) SEQUENCE DESCRIPTION: SEQ ID NO:22:
	420 421	CCACTAGTAC GCAAGTTCAC G 21
	1151	(2) INFORMATION FOR SEQ ID NO:47:
	1152	
	1153	(i) SEQUENCE CHARACTERISTICS:
	1154	(A) LENGTH: 109 amino acids
>	1155	(B) TYPE (PRT) amino our
	1156	(D) TOPOLOGY: Linear
	1157	
	1158	(xi) SEQUENCE DESCRIPTION: SEQ ID NO:47:
	1159	
	1160	Asp Ile Gln Met Thr Gln Ser Pro Ser Ser Leu Ser Ala Ser Val
	1161	1 5 10 15
	1162	
	1163	Gly Asp Arg Val Thr Ile Thr Cys Arg Ala Ser Lys Thr Ile Ser
	1164	20 25 30
	1165	
	1166	Lys Tyr Leu Ala Trp Tyr Gln Gln Lys Pro Gly Lys Ala Pro Lys
	1167	35 40 45
	1168	
	1169	Leu Leu Ile Tyr Tyr Ser Gly Ser Thr Leu Glu Ser Gly Val Pro

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RAW SEQUENCE LISTING PATENT APPLICATION US/09/234,182

DATE: 01/29/1999 TIME: 13:42:11

		INPUT SET: S30387.raw
	1170	50 55 60
	1171	
	1172	Ser Arg Phe Ser Gly Ser Gly Thr Asp Phe Thr Leu Thr
	1173	65 70 75
	1174	
	1175	Ile Ser Ser Leu Gln Pro Glu Asp Phe Ala Thr Tyr Tyr Cys Gln
	1176	80 85 90
	1177	
	1178	Gln His Asn Glu Tyr Pro Leu Thr Phe Gly Gln Gly Thr Lys Val
	1179	95 100 105
	1180	
	1181	Glu Ile Lys Arg
	1182	100 My hunth the arrest of the
	1183	Glu Ile Lys Arg 100 mly number the arrest outs when every 5 arrest a
	1250	(2) INFORMATION FOR SEQ ID NO:50:
	1251	
	1252	(i) SEQUENCE CHARACTERISTICS:
	1253	(A) LENGTH: 116 amino acids
>	1254	(B) TYPE: PRT
	1255	(D) TOPOLOGY: Linear
	1256	
	1257	(xi) SEQUENCE DESCRIPTION: SEQ ID NO:50:
	1258	
	1259	Glu Val Gln Leu Val Glu Ser Gly Gly Gly Leu Val Gln Pro Gly
	1260	1 5 10 15
	1261	
	1262	Gly Ser Leu Arg Leu Ser Cys Ala Ala Ser Gly Phe Ser Phe Thr
	1263	20 25 30
	1264	
	1265	Gly His Trp Met Asn Trp Val Arg Gln Ala Pro Gly Lys Gly Leu
	1266	35 40 45
	1267	
	1268	Glu Trp Val Gly Met Ile His Pro Ser Asp Ser Glu Thr Arg Tyr
	1269	50 55 60
	1270	
	1271	Ala Asp Ser Val Lys Gly Arg Phe Thr Ile Ser Arg Asp Asn Ser
	1272	65 70 75
	1273	
	1274	Lys Asn Thr Leu Tyr Leu Gln Met Asn Ser Leu Arg Ala Glu Asp
	1275	80 85 90
	1276	
	1277	Thr Ala Val Tyr Tyr Cys Ala Ala Arg Gly Ile Tyr Phe Tyr Gly
	1278	95 100 105
	1279	
	1280	Thr Thr Tyr Phe Asp Tyr Trp Gly Gln Gly Thr
	1281	110 115 116
	1282	
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